

Altered Differentiation of Hepatocytes in a Transgenic Mouse Model of Hepatocarcinogenesis*

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ABSTRACT

Transgenic mice carrying the SV40 T antigen (TA_g) gene, which develop hepatocellular and biliary cell tumors by 4 mo of age, show ductular structures in the neonatal liver. Coexpression of *c-myc* with TA_g increases the extent and persistence of ductular lesions and also accelerates tumor development. To analyze possible links between altered gene expression and cell differentiation and to determine the relationship between the ductular structures and tumor development in these mice, ductular cells in single (TA_g) and bitransgenic (TA_g × *c-myc*) mice were characterized for biliary and hepatocellular differentiation, transgene expression, and proliferation activity. The results show that the ductular cells in these transgenic mice have characteristics of biliary cells, including basement membrane formation, positive laminin staining, and bile duct-specific lectin (*Dolichos biflorus* agglutinin and peanut agglutinin) binding, and characteristics of hepatocytes, including albumin expression and ultrastructural features such as round nuclei with 1 or 2 nucleoli and well-developed cytoplasmic organelles. However, differences in transgene expression and cell proliferation between the ductular cells and nonductular hepatocytes were not apparent. Thus, the ductular cells could not be defined as tumor progenitor cells in these mouse livers. However, this model suggests that manipulation of gene expression can alter differentiation of hepatic parenchymal cells.

Keywords. Liver; neoplasms; ductular hepatocytes; cell differentiation; cell proliferation; bitransgenic mice

INTRODUCTION

Hepatocytes have been shown to form ductlike structures in some diseases. In humans, ductular hepatocytes have been reported in cases of alcoholic and cholestatic liver diseases, fulminant hepatitis, liver allograft rejection, and massive hepatic necrosis (17, 27, 30–32). In rats, ductular cells have been identified following treatment with chemicals (e.g., furan, D-galactosamine, carbon tetrachloride, and N-2-acetylaminofluorene in a choline-deficient diet) with or without bile duct ligation (5, 7, 14, 25, 26). In most instances, either extensive or chronic loss of hepatic parenchyma appears to be a necessary requisite for the appearance of ductular structures. In many of these cases, the ductular cells show characteristics of both hepatocytes and biliary cells and have sometimes been referred to as intermediate cells (5, 14, 17, 27, 30).

Transgenic mice carrying the simian virus 40 (SV40) T antigen (TA_g) gene fused to an albumin promoter/enhancer develop hepatocellular and biliary cell tumors at 3 or 4 mo of age (22). Examination of livers from these mice revealed the presence of ductular structures in neonates at 1 and 2 wk of age (8). Coexpression of a second transgene, *c-myc*, with TA_g accelerated tumor development dramatically (22) and also resulted in an increase in the extent and persistence of the ductular structure formation (8). In these bitransgenic mice, some hepatocellular and biliary cell tumors coexisted with the ductular

cells. Such phenotypic changes in hepatocytes prior to and/or during initial phases of tumor development led to speculation that the duct formation might be a step in the carcinogenic process underlying some tumors in this hepatocarcinogenesis model.

Histomorphologic features of a cell reflect differential gene expression. The effects of altered gene expression, however, are not necessarily limited to the change in morphology but may also involve other fundamental properties of the cell, such as growth and differentiation. The purposes of this study were to (a) analyze the possible links between altered gene expression and cell differentiation and (b) determine the relationship between the appearance of ductular hepatocytes and the development of neoplasia in this hepatocarcinogenesis model.

MATERIALS AND METHODS

Transgenic Mice. Animals carrying a single transgene, i.e., SV40 TA_g or *c-myc* gene fused to an albumin promoter/enhancer (AL), or bitransgenes, i.e., AL-TA_g × AL-*myc*, were used for the study. Details of the phenotype of each line of mouse have been described (8, 22). Transgene-negative littermates were used as controls.

Histopathology. Animals carrying single transgenes or bitransgenes were euthanatized at 1, 2, 3, or 4 wk of age by cervical dislocation or CO₂ asphyxiation (4–24 mice/line/age, see Table I), and livers were removed and fixed in 10% neutral buffered formalin or 4% paraformaldehyde. A part of each tissue was embedded in paraffin, sectioned, stained with hematoxylin and eosin (H&E) according to standard methods, and examined microscopically.

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TABLE I.—Incidence^a of ductular structures in transgenic mouse livers.

Mouse age (wk)	Transgenes			
	AL-TAg	AL-TAg × AL-myc	AL-myc	None
1	2/5 ^a (40)	7/9 (78)	0/7	0/8
2	9/24 (38)	17/18 (94)	0/11	0/17
3	0/7	8/9 (89)	0/10	0/5
4	0/12	0/7	0/4	0/5

^a Number of animals with ductular structures/number of animals examined (%).

Electron Microscopy. Liver tissues were sampled from animals carrying AL-TAg or AL-TAg × AL-myc at 7, 14, or 24 days of age (2 or 3 mice/line/age). A part of each liver was trimmed into small pieces and fixed in a glutaraldehyde/paraformaldehyde-mixed solution (2.6% glutaraldehyde, 2.0% paraformaldehyde, and 3.0% sucrose in 0.1 M phosphate buffer, pH 7.4) at 4°C. The tissues were processed according to routine methods, stained with uranyl acetate and lead citrate, and examined in a Zeiss EM 10C/CR electron microscope.

Lectin Histochemistry. Binding sites for 2 kinds of lectins, *Dolichos biflorus* agglutinin (DBA) and peanut agglutinin (PNA), were detected histochemically on formalin-fixed tissues from AL-TAg and AL-TAg × AL-myc mice at 1, 2, or 3 wk of age (2 mice/line/age). Deparaffinized sections were incubated in 3% hydrogen peroxide for 15 min and in phosphate-buffered saline containing 1% milk and 1% bovine serum albumin for 20 min, and then biotin-labeled lectins (Sigma Chemical Co., St. Louis, MO) were applied at a concentration of 30 µg/ml for 30 min followed by streptavidin-biotin complex (Biogenex Laboratories, San Ramon, CA) for 30 min. Signal was detected using diaminobenzidine (Sigma Chemical Co.) as the chromogen. The specificity of lectin binding was confirmed by preincubation of tissues with N-acetyl-D-galactosamine (Sigma Chemical Co.) or D(+)-galactose (Sigma Chemical Co.) before incubation with DBA or PNA, respectively.

Immunohistochemistry. Immunohistochemical staining for TAg was performed on formalin-fixed tissues from AL-TAg and AL-TAg × AL-myc mice at 1, 2, 3, or 4 wk of age (3 mice/line/age) as described previously (8). For detection of myc protein, sections of formalin-fixed, paraffin-embedded tissues from single transgenic and bi-transgenic mice at 1, 2, 3, or 4 wk of age (3 mice/line/age) were digested with 0.1% trypsin (Sigma Chemical Co.) in distilled water containing 0.1% calcium chloride for 20 min at 37°C and incubated in 3% hydrogen peroxide for 15 min. A rabbit polyclonal antibody to synthetic Myc peptide (OA-11-802, Cambridge Research Biochemicals, London, UK) was applied on the tissues at a dilution of 1:400 (2.5 µg/ml IgG) for 60 min at room temperature. The remainder of the staining was performed using Vectastain Elite ABC Kit (rabbit IgG, Vector Laboratories, Burlingame, CA). Sections of normal mouse colon were used as positive controls (16, 29). As a negative control, the primary antibody was replaced by normal rabbit IgG. Immunohistochemical staining for laminin was performed for the same animals as those used for lectin histochemistry. Formalin-fixed, paraffin-

embedded tissues were stained basically as described above except that protease (type XXIV, 4 mg/ml, Sigma Chemical Co.) and Vectastain ABC Kit (rabbit IgG) were used instead of trypsin and Vectastain Elite ABC Kit, respectively; the digestion by protease was performed for 15 min at room temperature. Primary antibody, a rabbit polyclonal anti-laminin antibody (Sigma Chemical Co.), was applied on the tissues at a dilution of 1:200 overnight at 4°C.

In situ Hybridization. A DNA probe of 668 bp, which overlaps the *HindIII* region of mouse albumin DNA used for *in situ* hybridization previously (2, 4, 12, 21), was generated using 2 rounds of polymerase chain reaction (PCR). Primers used were 5'-GTA TGC AGA AGT TTG GAG AGA-3' (upper) and 5'-CAC GAG AGT TGG GGT TGA CAC-3' (lower). Fifty microliters of reaction solution for the first round of PCR contained 0.4 µM of each primer, 8.2 µg of mouse liver cDNA library as a template, 0.2 mM of dNTPs, and 2 units of AmpliTaq polymerase (Perkin-Elmer, Foster City, CA) in 1× PCR buffer containing 1.5 mM MgCl₂. After incubation for 5 min at 94°C, Taq polymerase was added, and 35 cycles of PCR were performed under the following conditions: 1 min at 94°C, 1.5 min at 55 °C, and 1.5 min at 72°C, followed by 7 min at 72°C at the end of the last cycle for extension. The product was purified using a QIAquick Gel Extraction Kit (QUIAGEN, Chatsworth, CA), and the sequence was confirmed using an automated DNA sequencer (model 373A, Applied Biosystems, Foster City, CA). A second round of PCR was performed under the same conditions as the first round except that 150 ng of the purified product of the first round of PCR was used as a template, 0.07 mM of dUTP was replaced by digoxigenin (DIG)-11-dUTP (Boehringer-Mannheim, Indianapolis, IN), the cycle number was 10–30, and the final extension step was omitted. The DIG-labeled PCR product was used as a probe for *in situ* hybridization without further purification. As a negative control, the same size (668 bp) of probe for the murine heat-shock protein 70.2 gene (gift from Dr. Gotoh), which is expressed exclusively in the adult mouse testis (19), was used.

For *in situ* hybridization, sections of formalin-fixed, paraffin-embedded tissues from AL-TAg and AL-TAg × AL-myc mice at 1, 2, or 3 wk of age (2 mice/line/age) were digested with proteinase K at concentrations of 10–100 µg/ml; the temperature and the time for digestion were changed depending on the age of animals because tissues became more resistant to digestion with age. After digestion, the sections were refixed with 4% paraformaldehyde for 5 min and incubated with a prehybridization solution containing 50% deionized formamide and 0.3 M NaCl for ≥10 min, and then 20 µl of hybridization solution containing 5 ng/µl of probe DNA, 0.3 M NaCl, 50% deionized formamide, 1× Denhardt's solution, 20 mM Tris (pH 8.0), 5 mM EDTA, 10% dextran sulfate, 500 ng/µl tRNA, and 500 ng/µl salmon sperm DNA was applied. The slides with glass coverslips were incubated in a humidified chamber for 10–15 min at 100 °C to denature both probe DNA and tissue mRNAs and then kept overnight at 50°C. After hybridization, the tissues were rinsed in 2× standard saline citrate (SSC) for 3 min at

TABLE II.—Labeling index (LI) for livers of single transgenic and bitransgenic mice treated with BrdU at 14 days of age. Statistical analyses were not performed because of small group sizes.

Time after BrdU administration	Transgenes			
	AL-TAg	AL-TAg × AL-myc	AL-myc	None
2 hr				
n	3	6	5	2
LI (%) ^a				
Centrilobular	17.8 ± 1.1	25.6 ± 2.2	4.9 ± 1.7	4.4
Periportal	11.8 ± 2.2	15.5 ± 3.2	3.7 ± 1.3	3.6
Whole lobule	14.9 ± 1.7	20.8 ± 1.4	4.3 ± 1.4	3.9
3 days				
n	5	5	9	5
LI	ND ^b	ND	ND	ND
7 days				
n	12	2	8	6
LI	ND	ND	ND	ND

^a For n ≥ 3, mean ± SD; for n = 2, mean only.

^b ND = not determined.

room temperature and washed twice in 0.1× SSC for 1 hr at 60°C. Signal was detected using alkaline phosphatase-conjugated anti-DIG (Boehringer Mannheim) with nitroblue tetrazolium/5-bromo-4-chloro-3-indolyl phosphate as the chromogen. Tissues were counterstained with 0.02% fast green FCF and Gill's hematoxylin and mounted with aqueous mounting medium.

Cell Proliferation Analysis. Bromodeoxyuridine (BrdU, 30 mg/kg, ip) was administered to 2-wk-old mice carrying single transgenes or bitransgenes, and the animals were euthanatized 2 hr, 3 days, or 7 days after BrdU administration (2–12 mice/line/age, see Table II). Liver tissues were removed and fixed in 4% paraformaldehyde, and BrdU incorporation was detected immunohistochemically as described previously (28) using rat monoclonal BrdU antibody, MAS250 (Accurate Chemical and Scientific Corp., Westbury, NY). For each liver from animals euthanatized 2 hr after BrdU administration, a BrdU labeling index (LI = number of BrdU positive nuclei/total number of nuclei × 100) was obtained by counting ≥1,000 hepatocytes in the microscopic fields consisting of a minimum 5 centrilobular and 5 periportal areas selected at random. For those euthanatized 3 or 7 days after BrdU administration, distribution pattern of positive BrdU staining was examined microscopically, but LIs were not obtained.

RESULTS

Histopathology

Ductlike structures were found in mice carrying AL-TAg alone and in those carrying AL-TAg × AL-myc (Table I). The incidence, distinctiveness of the morphological features, and temporal persistence of these ductular structures were different between the single transgenic and bitransgenic mice.

In mice carrying AL-TAg alone, multiple small spaces, consistent with bile canaliculi and delimited by contiguous hepatocytes, were present at the periphery of liver lobes at 1 wk of age (Fig. 1). At 2 wk of age, the hepatocytes surrounding these spaces had a more discrete ductular pattern (Fig. 2). Ductular structures were seen also in centrilobular areas. At 3 wk of age, however, liv-

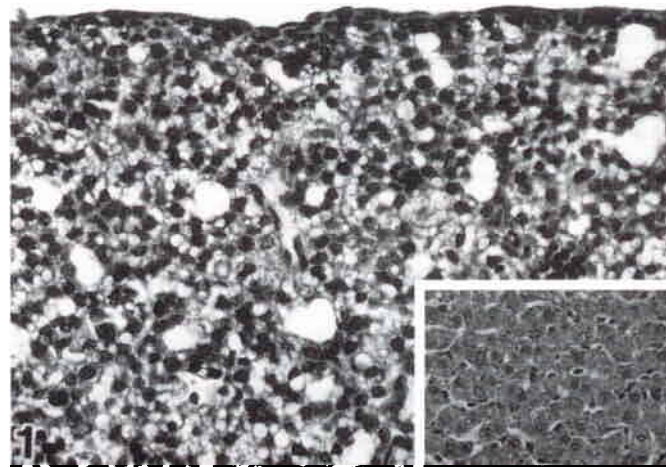


FIG. 1.—Multiple small spaces at the periphery of the liver lobe from a 1-wk-old AL-TAg mouse. H&E. ×340. Inset. Normal liver from a transgene-negative littermate of the same age. H&E. ×170.

ers were occupied by contiguous hepatocytes with varying degrees of dysplasia but without discrete hepatic cords, and ductular hepatocytes were no longer present. Focal proliferative lesions of biliary cells and hepatocytes were first seen at 4 and 6 wk, respectively, within the background of dysplastic hepatocytes; biliary and hepatocellular tumors were first observed at 6 and 8 wk, respectively (8).

In contrast to the mice carrying AL-TAg alone, mice coexpressing AL-myc with AL-TAg had more distinct ductular structures lined by hepatocytes, and these occupied a larger proportion of the liver and in centrilobular areas (Fig. 3); in some cases, entire liver lobules were occupied by the ductular structures. The cells lining the ductules were more basophilic than were nonductular he-

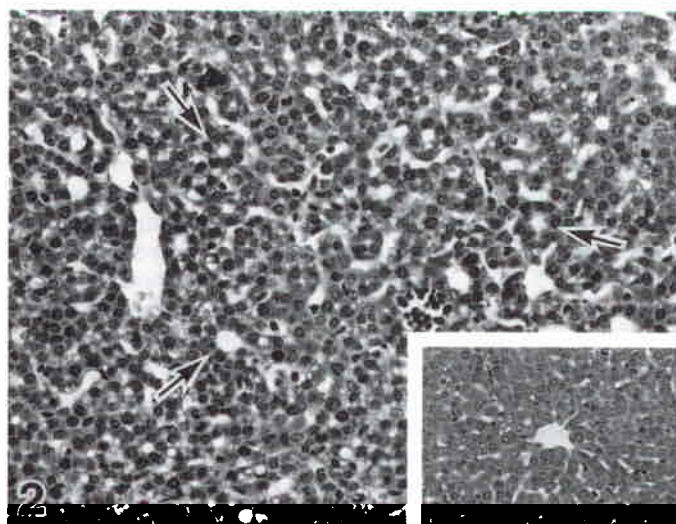


FIG. 2.—Ductular arrangement of some hepatocytes (arrows) in the liver from a 2-wk-old AL-TAg mouse. H&E. ×270. Inset. Normal liver from a transgene-negative littermate of the same age. H&E. ×140.

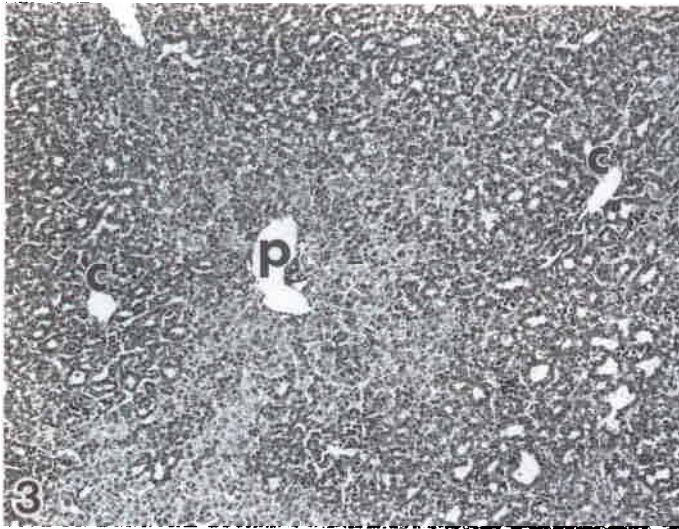


FIG. 3.—Centrilobular distribution of ductular hepatocytes in the liver from a 2-wk-old AL-TAg \times AL-myc mouse. C = central vein; P = portal vein. H&E. $\times 90$.

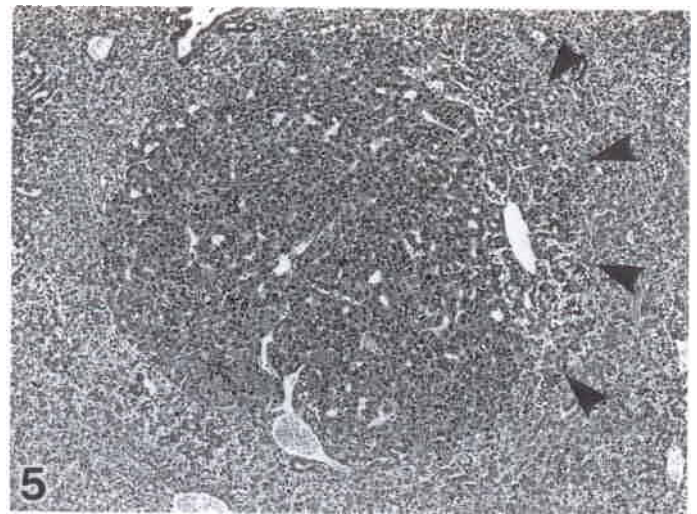


FIG. 5.—Focal proliferation of hepatocytes contiguous to a ductular area (arrowheads) in the liver from a 3-wk-old AL-TAg \times AL-myc mouse. H&E. $\times 85$.

patocytes but still had morphological features of hepatocytes in H&E-stained sections (Fig. 4). Unlike the ductular structures in single transgenic AL-TAg mice, the ductular lesions were still present in 8 of 9 bitransgenic mice examined at 3 wk of age. Furthermore, proliferative focal lesions of biliary cells were first seen at 1 wk of age and those of hepatocytes at 2 wk in the bitransgenic mice; cholangiolar and hepatocellular tumors developed at 3 wk of age (8). Some of these lesions were located in or were contiguous with the ductular areas (Figs. 5 and 6). Biliary cells in the proliferative lesions were columnar, had hyperchromatic, elongated nuclei with a high nucleus/cytoplasm ratio, and were distinct from the cells forming the ductular structures.

Severe hepatic damage was not seen in either line of mouse although apoptotic cells/bodies, which became ap-

parent at 2 wk of age, increased in number with age in both lines. There were no ductular lesions in livers from mice carrying AL-myc alone or in those from transgene-negative littermates at any age examined (Table I).

Ultrastructural Characteristics

Cells forming ductular structures had ultrastructural characteristics of both hepatocytes and of biliary cells. In normal livers from transgene-negative littermates, bile canaliculi between contiguous hepatocytes had a small lumen filled with microvilli (Fig. 7). The lumen of ductular structures in transgenic mouse liver was distended, irregular in shape, contained scanty microvilli, and was surrounded by 3 to >10 cells (Figs. 8 and 9). In some instances, basement membrane was detected between duct-forming cells and adjacent nonductular hepatocytes

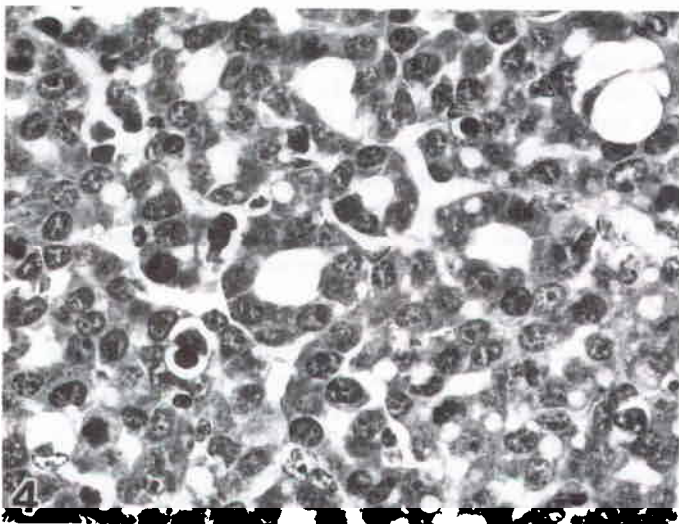


FIG. 4.—Higher magnification of ductular cells from the same liver as shown in Fig. 3. H&E. $\times 680$.

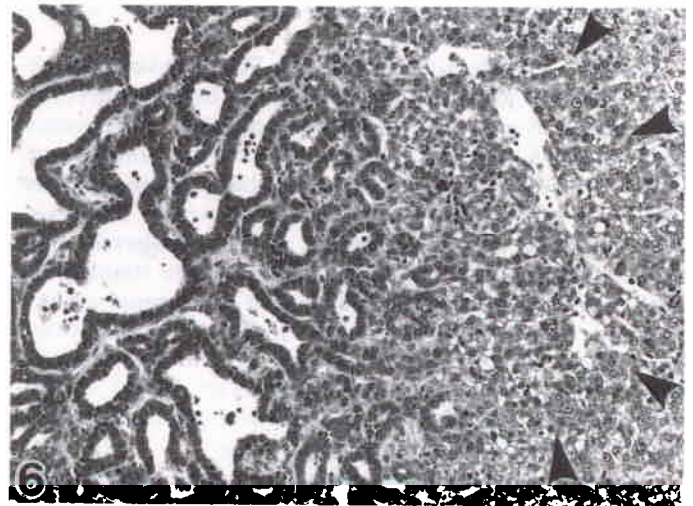


FIG. 6.—A biliary cell tumor contiguous to a ductular area (arrowheads) in the liver from a 3-wk-old AL-TAg \times AL-myc mouse. H&E. $\times 170$.

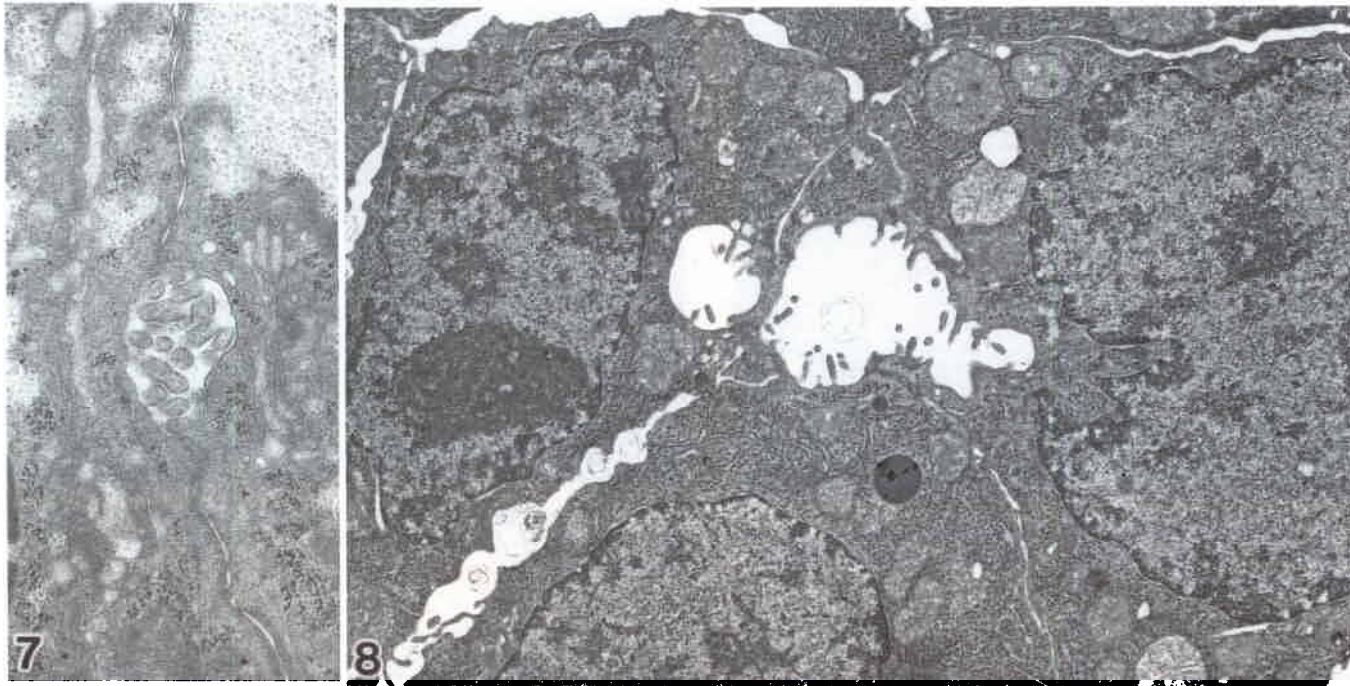


FIG. 7.—Electron micrograph of a normal bile canaliculus from a 2-wk-old nontransgenic littermate. The canaliculus has a small lumen of regular form filled with microvilli. Uranyl acetate and lead citrate. $\times 21,000$.

FIG. 8.—Electron micrograph of an early ductular structure from a 1-wk-old AL-Tag \times AL-myc mouse. Bile canaliculus is distended and contains scanty microvilli. Uranyl acetate and lead citrate. $\times 8,400$.

(Fig. 10). The cells forming the ductules, however, also had hepatocytic features, including centrally placed round nuclei with 1 or 2 nucleoli, numerous round mitochondria, well-developed rough endoplasmic reticulum and free ribosomes, and occasional lysosomes (Figs. 8 and 9).

Histochemical Characterization

Morphological and/or functional markers of biliary cells (bile duct-specific lectin binding and laminin) and of hepatocytes (albumin) were detected as follows in cells forming ductular structures.

Lectins. Two lectins, DBA and PNA, have been reported to bind to biliary cells but not to hepatocytes (33). In normal livers from transgene-negative littermates, DBA or PNA binding was detected on the luminal surface of bile ducts and/or in the cytoplasm of biliary cells; there was no positive staining in hepatocytes. However, in livers from single transgenic and bitransgenic mice carrying AL-Tag, weak but clearly positive staining for DBA and PNA was detected either on the luminal surface or in the cytoplasm of some cells forming ductular structures (Figs. 11 and 12). No binding sites for these lectins were detected in nonductular hepatocytes. There was no recognizable difference in the staining pattern between single transgenic and bitransgenic mice.

Laminin. In livers from single transgenic and bitransgenic mice carrying AL-Tag, laminin was detected around each ductular structure (Fig. 13) and around normal bile ducts. Increased staining intensity was associated with the distinctiveness of ductular structures. There was

no staining for laminin around hepatocytes that were not forming ducts. When ductular structures were not detected in the AL-Tag single transgenic mice at 3 wk of age, laminin was undetectable except for that surrounding normal bile ducts, whereas it remained positive around the more persistent ductular structures in the livers from bitransgenic mice. Positive staining for laminin was clearly seen around hyperplastic and neoplastic bile ducts in older AL-Tag and bitransgenic mice.

Albumin. In livers from nontransgenic littermates, albumin mRNA was detected in almost all hepatocytes, with highest staining intensity in periportal areas; albumin mRNA was not detected in normal biliary cells under the staining conditions. In livers from the AL-Tag and bitransgenic mice, cells forming ductular structures and nonductular hepatocytes were both albumin positive; even in cases where the ductular lesions occupied entire lobules, cells forming the ductules expressed albumin mRNA whereas biliary cells in the portal areas of the same sections were albumin negative (Fig. 14).

Transgene Expression

Tag. The staining intensity, pattern, and duration of TAG expression were the same between the single transgenic and bitransgenic mice carrying AL-Tag. TAG protein was present in the nuclei of most hepatocytes in 1-wk-old mice, including those hepatocytes surrounding the small spaces at the periphery of liver lobes (the earliest change of ductular structure formation). TAG-positive cells, however, gradually decreased with age. At 2 wk, TAG protein was detected mainly in centrilobular areas

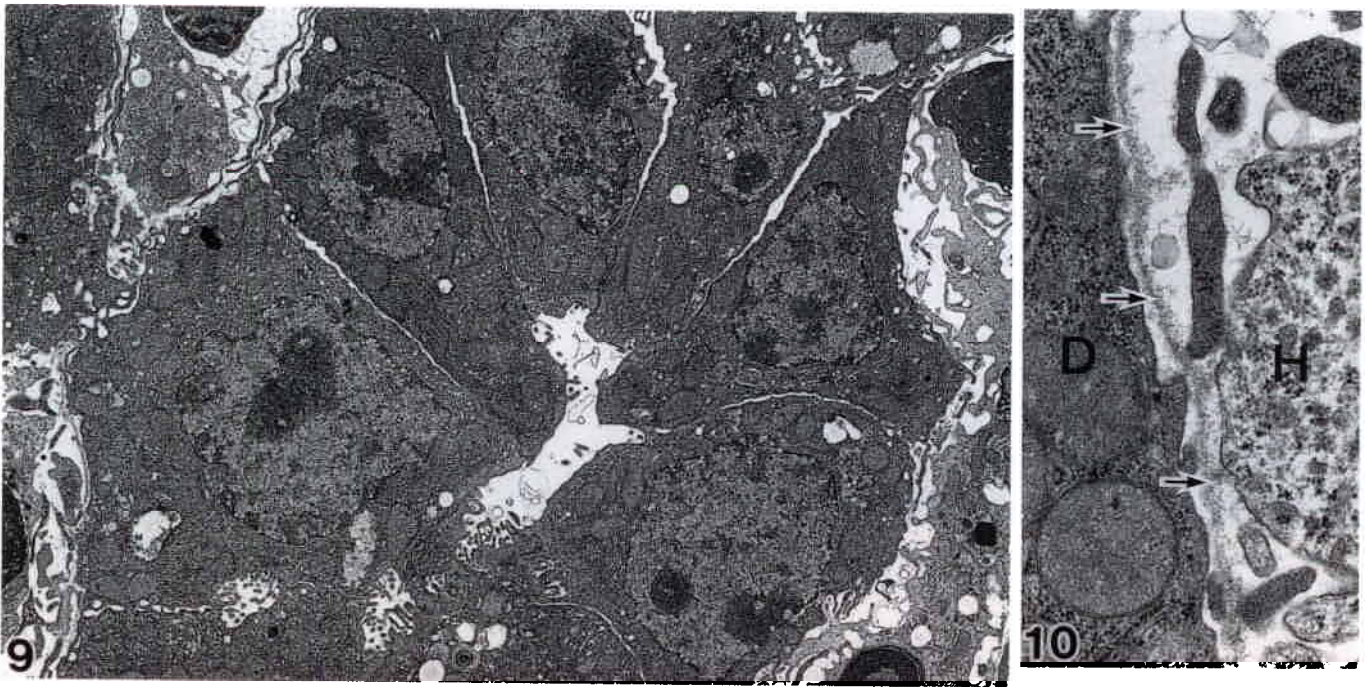


FIG. 9.—Electron micrograph of a ductular structure surrounded by several cells in the liver from a 1-wk-old AL-TAg \times AL-myc mouse. Uranyl acetate and lead citrate. $\times 4,500$.

FIG. 10.—Electron micrograph of a basement membrane (arrows) between a ductular cell (D) and nonductular hepatocyte (H). Uranyl acetate and lead citrate. $\times 27,500$.

(Fig. 15) where, in some livers, ductular structures were present. There was, however, no clear difference in TAg expression between cells forming ductular structures and nonductular hepatocytes. By 3 wk, most hepatocytes were TAg negative except for a small number of cells around central veins. TAg-positive biliary cells were also seen in some cases.

Myc. Myc protein was detected in the nuclei of most hepatocytes at 1–4 wk of age in both nontransgenic and transgenic mouse livers (Fig. 16). No detectable difference in myc expression was present among the livers from animals carrying AL-myc or AL-TAg \times AL-myc and transgene-negative littermates. Differences in myc expression between cells forming ductular structures and nonductular hepatocytes were not apparent.

Cell Proliferation Activity

The greatest degree of cell proliferation was seen in bitransgenic mice carrying AL-TAg \times AL-myc followed by AL-TAg single transgenic mice, as shown by BrdU LIs from those mice euthanatized 2 hr after BrdU administration (Table II). The LI in the centrilobular areas in bitransgenic mice was 1.7 times higher than that in the periportal areas; the same pattern of BrdU positivity was seen in livers from AL-TAg single transgenic mice. There were no recognizable differences in BrdU positivity between cells forming ductular structures and nonductular hepatocytes in single transgenic and bitransgenic mouse livers, although a BrdU LI was not obtained separately for these 2 types of cells because of the difficulty of distinguishing ductular and nonductular cells clearly in sections stained for BrdU. Active proliferation of cells in

bitransgenic mouse livers and, to a lesser extent, in the AL-TAg mouse livers was also indicated by apparent loss of BrdU-positive cells in their livers by 3 days after BrdU administration. In animals euthanatized 3 or 7 days after a pulse dose of BrdU, solid positive (brown) staining for BrdU was lost in many hepatocytes in single transgenic and bitransgenic mice carrying AL-TAg, and instead fine or coarse patches of brown staining were present in some hepatocellular nuclei, whereas persistent solid brown staining was detected in livers from AL-myc and transgene-negative littermates at the same time points.

DISCUSSION

We have demonstrated the transient occurrence of ductular structures in livers of mice carrying the AL-TAg transgene and shown that the ductular cells have morphological, histochemical, and functional characteristics of both hepatocytes and biliary cells. The ductular structures appeared to develop as dilated bile canaliculi between contiguous hepatocytes, and the cells forming these ductules exhibited evidence of biliary differentiation, including basement membrane, positive laminin staining, and bile duct-specific lectin binding, while maintaining features of hepatocytes, including albumin synthesis and ultrastructural characteristics such as round nuclei with 1 or 2 nucleoli and well-developed cytoplasmic organelles. Similar duct formation previously documented in humans (17, 27, 30–32) and animals (5, 7, 14, 25, 26) is typically associated with extensive or continuous loss of hepatic parenchyma. However, the livers of AL-TAg mice in the present study lacked evidence of massive hepatic cell loss, suggesting that this altered dif-

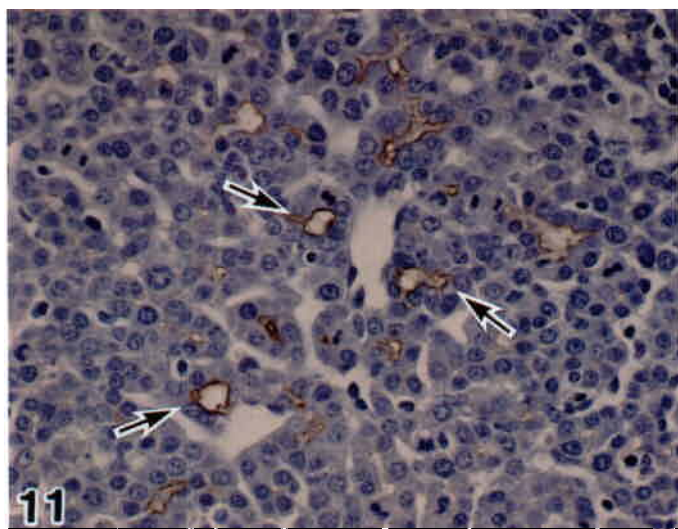


FIG. 11.—DBA binding on the luminal surface of ductular cells (arrows) in the liver from a 2-wk-old AL-TAg \times AL-myc mouse. Lectin histochemistry. $\times 370$.

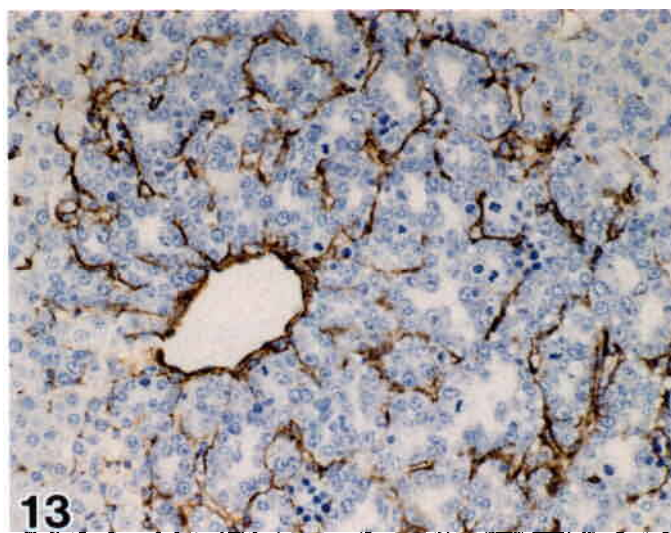


FIG. 13.—Laminin surrounding ductules in the liver from a 2-wk-old AL-TAg \times AL-myc mouse. Immunohistochemical staining for laminin. $\times 220$.

ferentiation is directly or indirectly attributed to the AL-TAg transgene and is not secondary to hepatic damage.

Whether TAg expression led to the altered differentiation by directly affecting cellular differentiation processes or by other indirect mechanisms is not known. The temporal and topographic occurrence of ductular structures appeared to be associated with TAg expression. Both ductular structures and TAg expression were detectable mainly at 1–2 wk of age; although TAg expression diminished with age, it remained longer in centrilobular areas than in periportal areas, the distribution pattern of which was consistent with that of ductular structures. However, ductular structures have not been reported in livers of other lines of mice carrying TAg (1,

3, 6, 11, 20, 23, 24) despite the fact that bile duct proliferation and/or biliary cell tumors have been reported in some of these mouse lines (1, 20, 23, 24). The early and transitory appearance of ductular structures might be one of the reasons for the absence of such reports. Another possibility is that the disrupted expression of some gene(s), caused by the insertion of the transgene (10, 18), but not the TAg gene itself, triggered the ductular formation in the present model. The present study could not reveal whether TAg itself was the causative factor for these phenotypic changes; however, this model appears to demonstrate that a genetic alteration introduced into the mouse genome could affect differentiation of hepatic cells.

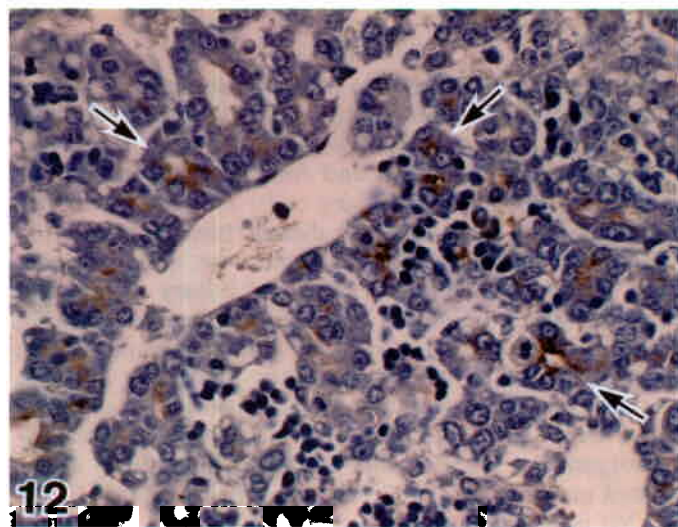


FIG. 12.—PNA binding on the luminal surface and in the cytoplasm of ductular cells (arrows) in the liver from a 2-wk-old AL-TAg \times AL-myc mouse. Lectin histochemistry. $\times 370$.

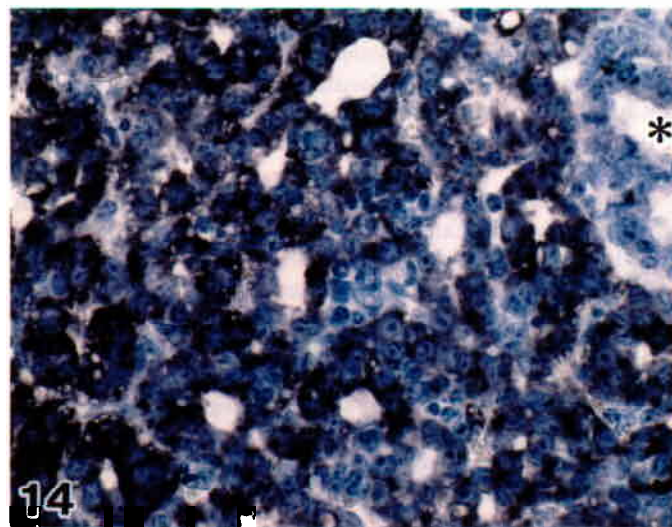


FIG. 14.—Albumin expression (black stain) in ductular cells in the liver from a 2-wk-old AL-TAg \times AL-myc mouse. Bile duct (*) cells are albumin negative. *In situ* hybridization for albumin mRNA. $\times 440$.

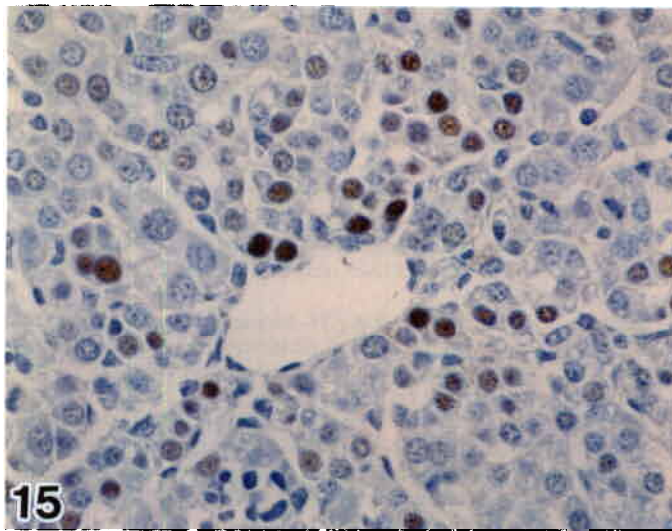


FIG. 15.—TAG expression in centrilobular hepatocytes in the liver from a 2-wk-old AL-TAG mouse. Immunohistochemical staining for TAG protein. $\times 440$.

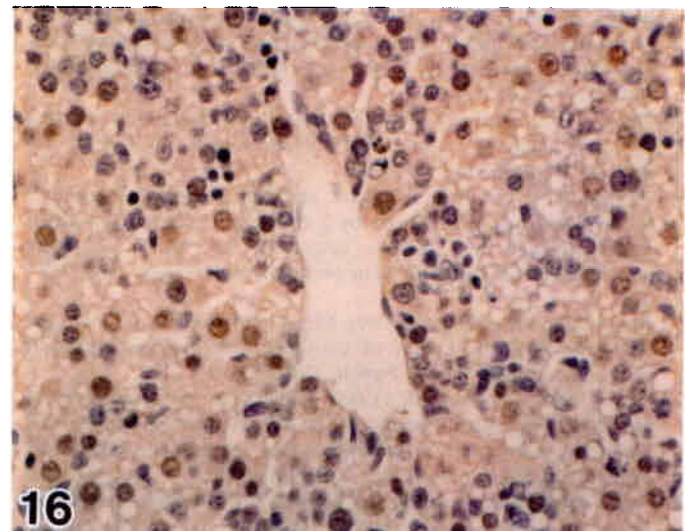


FIG. 16.—Myc expression in most hepatocytes in a 1-wk-old AL-myc mouse liver. Immunohistochemical staining for myc protein. $\times 370$.

Expression of the second transgene, AL-myc, caused an increase in the extent and persistence of the lesions possibly by a mechanism other than increasing the level of TAG expression. Continuous overexpression of *myc*, even at a level not detected by immunohistochemistry, also accelerated tumor development in AL-TAG mouse liver (8). In the present study, BrdU LIs in 2-wk-old AL-TAG \times AL-myc bitransgenic mice euthanatized 2 hr after BrdU dosing were generally higher than those in AL-TAG single transgenic mice. Although the number of mice examined is too small (3–6) to demonstrate differences statistically, the data suggest that coexpression of *c-myc* with TAG increased cell proliferation. The *c-myc* oncogene has been reported to be highly expressed in proliferating cells and downregulated by differentiation stimuli (13, 15). The present model may show that manipulation of *c-myc* expression could induce changes in both proliferation and differentiation of hepatic parenchymal cells *in vivo*.

In single transgenic and bitransgenic mice carrying TAG, high BrdU LIs (i.e., high cell proliferation activities) were observed in centrilobular areas. This activity may be related to the prolonged expression of TAG in these areas; TAG is known to have mitogenic activity (9). However, whether or not these high LIs were due to the presence of ductular cells could not be determined because LIs in ductular and nonductular cells could not be obtained separately; thus, the relationship between ductular structure formation and tumor development remained obscure. We also attempted to detect BrdU-positive cells in the livers 3 or 7 days after BrdU pulse dosing because we intended to use BrdU as a possible marker of ductular cells to trace their fate. However, both ductular and nonductular cells were actively proliferating in young (2-wk-old) mice, so that BrdU could not be used to differentially mark specific cell populations.

Despite the fact that duct formation occurs in some

hepatic diseases, little information is available on the molecular events associated with this phenotypic alteration or the relationship, if any, to neoplasia. We could not define a causative link between altered differentiation (ductular structure formation) and tumor development in this hepatocarcinogenesis model. Ductular formation may not have played a direct role in hepatic tumor development but rather may have reflected a perturbation in cellular differentiation that was indirectly linked to development of neoplasia. The present study does show, however, that a single genetic manipulation (the presence of a transgene) affects multiple genes governing morphological and functional characteristics of hepatocytes and biliary cells in a coordinated manner, which may indicate the existence of 1 or more key signal transduction pathways whose altered expression is responsible for the change in hepatic parenchymal cell differentiation. Elucidating the underlying molecular changes associated with altered differentiation states should contribute to our understanding of the mechanisms of differentiation of hepatic parenchymal cells.

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